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## RAW SEQUENCE LISTING

DATE: 06/27/2002

PATENT APPLICATION: US/09/998,904

TIME: 19:18:55

Input Set : N:\Crf3\06112002\I998904.raw

Output Set: N:\CRF3\06272002\I998904.raw

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1 <110> APPLICANT: The University of Texas Board of Regents
2 <120> TITLE OF INVENTION: Prediction of Disease-Causing Alleles from Sequence
3 Context
4 <130> FILE REFERENCE: 119929-1037
5 <140> CURRENT APPLICATION NUMBER: US/09/998,904
6 <141> CURRENT FILING DATE: 2002-06-04
7 <160> NUMBER OF SEQ ID NOS: 12
8 <170> SOFTWARE: PatentIn version 3.1
10 <210> SEQ ID NO: 1
11 <211> LENGTH: 101
12 <212> TYPE: DNA
13 <213> ORGANISM: mammal
14 <220> FEATURE:
15 <221> NAME/KEY: misc_feature
16 <222> LOCATION: (51)..(51)
17 <223> OTHER INFORMATION: n = a or g
18 <400> SEQUENCE: 1
W--> 19      gctggggccaa gctctacagc ttggtgatct ggggggtgtac gctgctcctg ngctcaccca      60
20      tgctggtgtt ccggaccatg aaggagtaca gcgatgaggg c      101
22 <210> SEQ ID NO: 2
23 <211> LENGTH: 101
24 <212> TYPE: DNA
25 <213> ORGANISM: mammal
26 <220> FEATURE:
27 <221> NAME/KEY: misc_feature
28 <222> LOCATION: (51)..(51)
29 <223> OTHER INFORMATION: n = c or g
30 <400> SEQUENCE: 2
W--> 31      gctgggtgttc cggaccatga aggagtacag cgatgagggc cacaacgtca ncgcttgtgt      60
32      catcagctac ccatecctca tctgggaagt gttcaccaac a      101
34 <210> SEQ ID NO: 3
35 <211> LENGTH: 101
36 <212> TYPE: DNA
37 <213> ORGANISM: mammal
38 <220> FEATURE:
39 <221> NAME/KEY: misc_feature
40 <222> LOCATION: (51)..(51)
41 <223> OTHER INFORMATION: n = a or g
42 <400> SEQUENCE: 3
W--> 43      gggccaagct ctacagcttg gtgatctggg ggtgtacgct gctcctgagc ncacccatgc      60
44      tgggtgttccg gaccatgaag gagtacagcg atgagggcca c      101
46 <210> SEQ ID NO: 4
47 <211> LENGTH: 101

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48 <212> TYPE: DNA
49 <213> ORGANISM: mammal
50 <220> FEATURE:
51 <221> NAME/KEY: misc_feature
52 <222> LOCATION: (51)..(51)
53 <223> OTHER INFORMATION: n = a or g
54 <400> SEQUENCE: 4
W--> 55      ctgcccttct gggccatcac catctccaac aacttcgact ggctcttttg ngagacgctc      60
56      tgccgcgtgg tgaatgcat tatctccatg aacctgtaca g      101
58 <210> SEQ ID NO: 5
59 <211> LENGTH: 101
60 <212> TYPE: DNA
61 <213> ORGANISM: mammal
62 <220> FEATURE:
63 <221> NAME/KEY: misc_feature
64 <222> LOCATION: (51)..(51)
65 <223> OTHER INFORMATION: n = c or t
66 <400> SEQUENCE: 5
W--> 67      cttctgggcc atcaccatct ccaacaactt cgactggctc tttggggaga ngctctgccg      60
68      cgtggtgaat gccattatct ccatgaacct gtacagcagc a      101
70 <210> SEQ ID NO: 6
71 <211> LENGTH: 101
72 <212> TYPE: DNA
73 <213> ORGANISM: mammal
74 <220> FEATURE:
75 <221> NAME/KEY: misc_feature
76 <222> LOCATION: (51)..(51)
77 <223> OTHER INFORMATION: n = g or a
78 <400> SEQUENCE: 6
W--> 79      ggacaccggc caccctccgc gccaccacc ctcgctttct ccggcttcct ntggcccagg      60
80      cgccgcgcgcg acccggcagc tgtctgcgca cgccgagctc c      101
82 <210> SEQ ID NO: 7
83 <211> LENGTH: 101
84 <212> TYPE: DNA
85 <213> ORGANISM: mammal
86 <220> FEATURE:
87 <221> NAME/KEY: misc_feature
88 <222> LOCATION: (51)..(51)
89 <223> OTHER INFORMATION: n = g or a
90 <400> SEQUENCE: 7
W--> 91      ctgtctcagc agtggacagc gggcatgggt ctgctgatgg cgctcatcgt nctgctcatc      60
92      gtggcgggca atgtgctggt gatcgtyggc atcgccaaga c      101
94 <210> SEQ ID NO: 8
95 <211> LENGTH: 101
96 <212> TYPE: DNA
97 <213> ORGANISM: mammal
98 <220> FEATURE:
99 <221> NAME/KEY: misc_feature
100 <222> LOCATION: (51)..(51)

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101 <223> OTHER INFORMATION: n = t or c
102 <400> SEQUENCE: 8
W--> 103      cgggcatggg tctgctgatg gcgctcatcg tgctgctcat cgtggcgggc natgtgctgg      60
104      tgatcgtggc catcgccaag acgccgcggc tgcagacgct c      101
106 <210> SEQ ID NO: 9
107 <211> LENGTH: 101
108 <212> TYPE: DNA
109 <213> ORGANISM: mammal
110 <220> FEATURE:
111 <221> NAME/KEY: misc_feature
112 <222> LOCATION: (51)..(51)
113 <223> OTHER INFORMATION: n = a or g
114 <400> SEQUENCE: 9
W--> 115      ctgctgatgg cgctcatcgt gctgctcatc gtggcgggca atgtgctggt natcgtggcc      60
116      atcgccaaga cgccgcggct gcagacgctc accaacctct t      101
118 <210> SEQ ID NO: 10
119 <211> LENGTH: 101
120 <212> TYPE: DNA
121 <213> ORGANISM: mammal
122 <220> FEATURE:
123 <221> NAME/KEY: misc_feature
124 <222> LOCATION: (51)..(51)
125 <223> OTHER INFORMATION: n = t or c
126 <400> SEQUENCE: 10
W--> 127      tcgccaagac gccgcggctg cagacgctca ccaacctctt catcatgtcc ntggccagcg      60
128      ccgacctggt catggggctg ctggtggtgc cgttcggggc c      101
130 <210> SEQ ID NO: 11
131 <211> LENGTH: 101
132 <212> TYPE: DNA
133 <213> ORGANISM: mammal
134 <220> FEATURE:
135 <221> NAME/KEY: misc_feature
136 <222> LOCATION: (51)..(51)
137 <223> OTHER INFORMATION: n = a or g
138 <400> SEQUENCE: 11
W--> 139      cgtggtgtgg ggccgctggg agtacggctc cttcttctgc gagctgtgga nctcagtggg      60
140      cgtgctgtgc gtgacggcca gcatcgagac cctgtgtgtc a      101
142 <210> SEQ ID NO: 12
143 <211> LENGTH: 100
144 <212> TYPE: DNA
145 <213> ORGANISM: mammal
146 <220> FEATURE:
147 <221> NAME/KEY: misc_feature
148 <222> LOCATION: (51)..(51)
149 <223> OTHER INFORMATION: n = t or c
150 <400> SEQUENCE: 12
151      ttccgctacc agagcctgct gacgcgcgcg cgggcgcggg gcctcgtgtg accgtgtggg      60
152      ccatctcggc cctggtgtcc ttctgccc tcctcatgca      100

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RAW SEQUENCE LISTING ERROR SUMMARY  
PATENT APPLICATION: US/09/998,904

DATE: 06/27/2002  
TIME: 19:18:56

Input Set : N:\Crf3\06112002\I998904.raw  
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:1; N Pos. 51  
Seq#:2; N Pos. 51  
Seq#:3; N Pos. 51  
Seq#:4; N Pos. 51  
Seq#:5; N Pos. 51  
Seq#:6; N Pos. 51  
Seq#:7; N Pos. 51  
Seq#:8; N Pos. 51  
Seq#:9; N Pos. 51  
Seq#:10; N Pos. 51  
Seq#:11; N Pos. 51